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FIG. 2

SEQ ID NO: 4	1	60
SEQ ID NO: 2	KIGRRNAA....	
SEQ ID NO: 5	AGQISGMD.EPLEI	
		MANQDLIPVVVNGAAGKMGREVIKAVAQAPDLQLVGAVDHNPSLQGGDIGEVVGIAPLEV	
SEQ ID NO: 4	61KVLCSQTQMPSPQSTI.....	120
SEQ ID NO: 2	KVIIGATKEIGRTAIAAVSKARGMELAGAIID.	
SEQ ID NO: 5		PVLNDLTMVLGSLAQSRATGVVDFSEPSAVYDNVKQAAAGLSSVWYVPKIELETVTTEL	
		PVLADLQSVLVLATQEKIQGVMDFTHPSGVYDNVRSIAIAYGVRPVVGTTLGLSEQQIQDL	
SEQ ID NO: 4	121S.QCI...GLDAGEI.....	180
SEQ ID NO: 2		SAFCEKAS.GCLVAPTLSIGSVLLQQAIIQASFHYSNVEIVESRPNP.SDLPQDRIQIA	
SEQ ID NO: 5		GDFAEKASTGCLIAPNFAIGVLLMQQAQAVQACQYFDHVEIIELHHNQKADAPSGTAIKTA	
SEQ ID NO: 4	181	..LNDLTMV.....LGSIAQTRA.....	240
SEQ ID NO: 2		NNISDLGQIYNR...EDMDSSSPARGQLLGEDGVRVHSMVLPGLVSSSTINFSGGPGMYT	
SEQ ID NO: 5		QMLAEMGKTENPPAVEEKETIAGAKGGL.GPGQIPIHISIRLPGLIAHQEVLFSGSPGQLYT	
SEQ ID NO: 4	241	NVKQA.....	276
SEQ ID NO: 2		LRHDVANVQCLMPGLILAIRKVVRFKNLIYGLEKFL	
SEQ ID NO: 5		IRHDTTDRACYPGVLGIRKRWELKGLVYGLEKLL	

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FIG. 3

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SEQ ID NO: 7 1 L..... 60
SEQ ID NO: 9 VS.....
SEQ ID NO: 11 MAITATISVPLTSPSRRTLTSVNSLSPLSTRSLTPQRTFKYPNSRLVWSSMSTETAVK
SEQ ID NO: 13 .....
SEQ ID NO: 14 .....

SEQ ID NO: 7 61 ..... 120
SEQ ID NO: 9 .....
SEQ ID NO: 11 TSSASFLNRKESGFLHFAKYHGLGNDVFLIDNRDSSSEPKISAEKAVQLCDRNFVGADGV
SEQ ID NO: 13 .....ALHFVKYQGLGNDFIMVDNRDSAVPKVTPEEAAKLCDRNFGXGADGV
SEQ ID NO: 14 .....MALSFESKYHGLGNDFILVDNRQSTEPCLTPDQAQQLCDRHFEGIGADGV

SEQ ID NO: 7 121 ..... 180
SEQ ID NO: 9 IFVMPGVNGADYTMRIFNSDGSEPEMCGNGVRCFARFIAELENLQGTNRFTIHTGAGKIV
SEQ ID NO: 11 IFVLPGISGTDYTMRIFNSDGSEPEMCGNGVRCFARFIAELENLQGTSHFKIHTGAGLII
SEQ ID NO: 13 IFVLPGVNGADYTMRIFNSDGSNNRWX.GFV.....
SEQ ID NO: 14 IFALPGQGGTDYTMRIFNSDGSEPEMCGNGIRCLAKFLADLEGVEEK.TYRIHTLAGVIT

SEQ ID NO: 7 181 ..... 240
SEQ ID NO: 9 PEIQSDGQVKVDMGEPILSGLDIPTKLLATKNKAVVQAEALAVEGLTWHVTCVSMGNPHCV
SEQ ID NO: 11 PEVLEDGNVRVDMGEPVLKALDVPTKLPAKNDNAVVKSQLVVDGVIWHVTCVSMGNPHCV
SEQ ID NO: 13 .....
SEQ ID NO: 14 PQLLADGQVKVDMGEPQLLAELIPTTLAPAGEK.VVDLPLAVAGQTWAVTCVSMGNPHCL
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FIG.3 (Continued)

SEQ ID NO: 7	241	TFGANELKVLQVDDLKLS	300
SEQ ID NO: 9		
SEQ ID NO: 11		TFSREGSQNLLVDELKLA	
SEQ ID NO: 13		
SEQ ID NO: 14		TFVDD.....VDSLNL	
		TEIGPLFEHHHPQFSQRT	
		NTFEFIQVLGSDRLKMR	
		VWERGAGITL	
SEQ ID NO: 7	301	ACGTGACAVVVA	359
SEQ ID NO: 9		
SEQ ID NO: 11		ACGTGACATVVA	
SEQ ID NO: 13		
SEQ ID NO: 14		ACGTGACATVVA	
		AVLTGRGDRRCTVEL	
		PGGNLEIWSAQDNRL	
		YMTGPAQRFVSGQAEI	

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FIG. 4

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1  ASSSLFQSLPESLQTSK.PYAPPKPAAHFVRA.....QSPLTQNNSSSKHRRPAD
50
SEQ ID NO:24
SEQ ID NO:25
SEQ ID NO:27
LSSCLFNASVSSLNPKQDPIRRHRSTSLLRHPVVISCTADGNNIKAPIETAVKPPHRT
E

61
SEQ ID NO:20
SEQ ID NO:22
SEQ ID NO:24
SEQ ID NO:26
SEQ ID NO:27
DNIRDEARR.NRSNAVNPFESAKYVPENA...APGSTESYSLSLDEIVYRSRSGGLLDVEHDM
120
.....MENGAATNGASEKSHSPS
ENIRDEARRINAPHDHHLFSAKYVPFNADSSSSSTESYSLSLDEIVYRSQSGGLLDVQHDM
.....
121
SEQ ID NO:16
SEQ ID NO:18
SEQ ID NO:20
SEQ ID NO:22
SEQ ID NO:24
SEQ ID NO:26
SEQ ID NO:27
EALKRFDGAYWRDLFDSRVGKSTWPYGSVWSKKKEWVLPEIDDDDDIVSAFEKNSNLFWAE
180
.....
QTYLSTRGDCYGLSFETV.....
DALKREDGEYWRNLFDSDRVGKTTWPYGSVWSKKKEWVLPEIHDDDDIVSAFEKNSNLFWAE
.....
181
SEQ ID NO:16
SEQ ID NO:18
SEQ ID NO:20
SEQ ID NO:22
SEQ ID NO:24
SEQ ID NO:26
SEQ ID NO:27
RFGKQFLGMNDLWVKHGGISHTGSFKDLGMTVLVSQVNRLRKMKRPVVGVCSTGDTSA
240
.....VGCASTGDTSA
LKGLAADGGLFPEEVPAAATEWQSWKDLPTYELAVK.....
RFGKQFLGMNDLWVKHCGISHTGSFKDLGMTVLVSQVNRLRKMKRPVVGVCSTGDTSA
.....
RFGKQFLGMNDLWVKHGGISHTGSFKDLGMTVLVSQVNRLRKMKRPVVGVCSTGDTSA

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FIG. 4 (Continued)

SEQ ID NO:16	241	ALSAYCAAAGIP	300
SEQ ID NO:18		ALFVLPADRLSLOQLIQPIANGATVLSLDTDFDGCMLIREVTAE	
SEQ ID NO:20		PI	
SEQ ID NO:22		LSLYISPAEVPTE	
SEQ ID NO:24		DLRALVER	
SEQ ID NO:26			
SEQ ID NO:27		ALSAYCASAAP	
		PSIVFLPANKISLAQLVQPIANGAFVLSIDTDFDGCMLIREVTAE	
		PI	
		LIQPIANGATVLSLDTDFDGCMLIREVTAE	
		PI	
		ALSAYCASAGIP	
		SIVFLPANKISMAQLVQPIANGAFVLSIDTDFDGCMLIREITAE	
		PI	
SEQ ID NO:16	301	YLANSLNPL	360
SEQ ID NO:18		RLEGQKTA	
SEQ ID NO:20		AEILQQFNWQVPDWVIVPGNGLNIYAFYKGFEMCRVLGLV	
SEQ ID NO:22			
SEQ ID NO:24		SYSTFRSKEVVPLVKLEDNHLLELFHGNYSF	
SEQ ID NO:26			
SEQ ID NO:27		YLANSLNSL	
		KLEGQKTA	
		AEILQQFDWQVPDWVIVPGSNGLNIYAFYKGFQELGLV	
		YLANSLNSL	
		XLEGQKTA	
		IRDIATXNQVPLGHIPRRQSTFYAF	
		LQGF	
		YLANSLNSL	
		RLEGQKTA	
		AEILQQFDWQVPDWVIVPGNGLNIYAFYKGFQELGLV	
SEQ ID NO:16	361	DRVPRLVCAQAANANPLYRY	420
SEQ ID NO:18		KSGWTEFEPQTAETTFASAIQIGDPVSVDR	
SEQ ID NO:20		VVALKATD	
SEQ ID NO:22			
SEQ ID NO:24		KDCALQFLGNLXEYF	
SEQ ID NO:26			
SEQ ID NO:27		DKIPRLVCAQAANADPLYLYFKSGWKEFKPVKSSTTFASAIQIGDPVSI	
		DRVHALKSCD	
		DRIPRMVCAQAANANPLYLHYKSGWKDFKPM	
		TASTTFASAIQIGDPVSI	
		DRVYALKKCN	
SEQ ID NO:16	421	GIVEEATEEELMDATALADRTGMFACPH	480
SEQ ID NO:18		TGVALAALFKLQQR	
SEQ ID NO:20		IIGPNDR	
SEQ ID NO:22			
SEQ ID NO:24		DAMVQADSTGMFICPHTGVALAALIKLRN	
SEQ ID NO:26		RGVIGAGERVV	
		STAHGL	
		GIVEEATEEELMDATAQADSTGMFICPHTG	
		VALTALFKLRNSGV	
		IKATDR	

FIG.4 (Continued)

SEQ ID NO: 27	GIVEEATEEEELMDAMAQADSTGMFICPHTGTGVALTALFKLRNQGVIAPTORTVVVSTAHGL
	481
SEQ ID NO: 16	KFTQSKIDYHDKNIKDMVCQYANPPISVKADEFGSVMVDVLQKN.....LNGKI...
SEQ ID NO: 18MACKYSNPPVSVKADEFGAVMDVLKKR.....LKGKL...
SEQ ID NO: 20
SEQ ID NO: 22	KFAQSKIDYHSGLIIPGMG.RYANPLVSVKADEFGSVMVDVLKDSCTTSPPTLTSLDVAK
SEQ ID NO: 24	KFTQSKIDYHDKDIKDMACRYANPPMQVKADEFGSVMVDVLKTY.....LQSKA..H
SEQ ID NO: 26
SEQ ID NO: 27	KFTQSKIDYHSNAIPDMACRFSNPPVDVKADEFGAVMDVLKSY.....LGSNTLTLS

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FIG. 5

SEQ ID NO: 29	1	60
SEQ ID NO: 31		
SEQ ID NO: 33		
SEQ ID NO: 34		MASHDYLLKKILTARVYDVAFETELEPARNLSARLRNPVYLKREDNQPVSEFKLRGAYNKM	
	61	120
SEQ ID NO: 29	T'V'LEGD	
SEQ ID NO: 31		
SEQ ID NO: 33		
SEQ ID NO: 34		AHIPADALARGVITASAGNHAQGVAFSAARMGVKAVIVVPVTPQVKVDVRAHGGSPGE	
	121	180
SEQ ID NO: 29		SYDEAQSAYAK.....LRCQQE.GRTFVPPFDHPDVTGQGTIGMEIVRQLQGFLHAI FVP	
SEQ ID NO: 31		
SEQ ID NO: 33		
SEQ ID NO: 34		VIQAGESYSDAYAHALKVQEEERGLTFVHPFDDPYVIAGQGTIAMEILRQHQQGPIHAI FVP	
	181	240
SEQ ID NO: 29		VGGGGLIAGIAAYVKVRPEVKIIGVEPSDANAMALSLCHGKRVMLEHVGGFADGVAVKA	
SEQ ID NO: 31		
SEQ ID NO: 33		
SEQ ID NO: 34		IGGGGLAAGVAAYVKVRPEIKVIGVQAEDESCAMAQSLQAGKRVELAEVGLFADGTAVKL	
	241	300
SEQ ID NO: 29		VGEETFRLCRELVDGIVMVS RDAICASI KOMFEKRSILEPAGALALAGAEAYCKYYNLK	
SEQ ID NO: 31		
SEQ ID NO: 33		
SEQ ID NO: 34		VGEETFRLCKEYLDGVVTVDTDALCAAIKDVFQDTRSVLEPSGALAVAGAKLYAERE GIE	

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FIG.5 (Continued)

301	360
SEQ ID NO:29	GETVVAITSGANMNFDRRLRVTELVGRKREAVLATFLPERQGSFKKFTLVGRMNITE
SEQ ID NO:31	..NIVAITSGANMNFDRRVVTELVANVGRKQEAVALTVMAEPPGSKQFCFCELVGQMNITE
SEQ ID NO:33
SEQ ID NO:34	NQTLVAVTSGANMNFDRMREFAERAEEVGEAREAVFAVTIPEERGSEKRFCSLVGDRNVTE
361	420
SEQ ID NO:29	FKYRYDSNAKDVLVSVGIYTDNELGAMMDRMESAKLRTVNLTNDLAKDHLRYFIGGR
SEQ ID NO:31	FKYRYNSNEK.AVVLYSVGVHTISELRAMQERMESQLKTYNLTESDLVKDHLRYLMGGR
SEQ ID NO:33
SEQ ID NO:34	FNRYRI.ADAQSAHIFVGVQIRRRGESADIAANFESHGFKTADLTHDELSKEHIRYMVGGR
421	480
SEQ ID NO:29	SEIK.DELVYRFIFPERPGALMKFLDTFSPRWNISLFHYRAQGEAGANVLVGIVPPAEF
SEQ ID NO:31	SNVQ.NEVVVSPXPRKTGALMKFLDXFSPRWDISL.....
SEQ ID NO:33RPGALMKFLDPSPRWNISLFHYRGEGETGANVLVGIVPKSEM
SEQ ID NO:34	SPLALDERLFRFEFFPERPGALMKFLSSMAPDWNISLFHYRNQGADYSSILVGLQVPQADH
481	512
SEQ ID NO:29	DEFKSHANNLGVEYMSEHNNNEIYRLLLRDPKV
SEQ ID NO:31
SEQ ID NO:33	DEFHDRANKLGVDYKVNNDDDDFQLLMH.....
SEQ ID NO:34	AEFERFLAALGYPYVEESANPAYRIFLS.....

SEQ	ID	NO:	36	1086	CCGAGCCTCTCTCCCGTGT	TTGTGACACAGTACGSCACCGGCGGATCCCGACAAAGGAGA	1145
SEQ	ID	NO:	37	1734	CAGAGCCACTGTCCGTAT	TCGTCGACACATACGGCACTGGCAGGATCCCTGACAAGGAGA	1793
SEQ	ID	NO:	36	1146	TCCTCAAGATTGTCAAGGAGAC	TTCGATTTTCAGGCCTGGCATGATTATCATCAACCTTG	1205
SEQ	ID	NO:	37	1794	TCCTCAAGATTGTGAAGGAGAACT	TCGACTTCAGGCCTGGCATGATCATCATCAACCTTG	1853
SEQ	ID	NO:	36	1206	ACCTCAAGAAAGCGCGCAACGSGCGCT	TACCTCAAGACGGCAGCCTACGGCCACTTCGGAA	1265
SEQ	ID	NO:	37	1854	ACCTCAAGAAAGCGCGCAACGACGCT	TACCTCAAGACGGCGGCTTACGGTCACTTCGGAA	1913
SEQ	ID	NO:	36	1266	GGGACGACCCCTGACTT	TCACCTGGGAGGTGTGAAGCCACTCAAGTCGGAGAAACCTTCTG	1325
SEQ	ID	NO:	37	1914	GGGACGACCCAGACTT	TCACCTGGGAGGTGTGAAGCCCTCAAGTGGGAGAGCCTTCTG	1973
SEQ	ID	NO:	36	1326	CCTAAGCGCGCCTTTT	1341	
SEQ	ID	NO:	37	1974	CCTAAAAGCTTCCCTTT	1989	

FIG. 7

[illegible]

[illegible]

[illegible]

[illegible]

SEQ	ID	No:	42	461	GCTATGCCACCGATGAGACCCCTGAGCTGATGCCCTCAAGCACGTGCTTGCCACCACAAGC	520
SEQ	ID	No:	43	443	GCTACGCCACTGATGAGACCCCTGAGCTGATGCCCTCACCCACATGCTTGGCACCAAGC	502
SEQ	ID	No:	42	521	TYGAGCTCGCCTCACSGAGGTCGCGAAGAATGGCACCTGCGCCTGGGTCAAGGCCTGACG	580
SEQ	ID	No:	42	503	TCGAGCTCGCCTCACCGAGGTCGCGAAGAATGGCACCTGCGCCTGGCTCAGGCCTGATG	562
SEQ	ID	No:	42	581	GAAGACCCAGGTCACAGTCGAGTACCTAAACGAGGATGGTGCCATGGTACCTGTTTCGTG	640
SEQ	ID	No:	43	563	GAAGACCCAGGTCACCATTGAGTACCTAAACGAGGGTGGTGCCATGGTGCCGTTTCGTG	622
SEQ	ID	No:	42	641	TGCACACCGTCCTCATCTCCACCCAGCAGCAGAGACCGTCAACCAACGACGAGATTGCTG	700
SEQ	ID	No:	43	623	TGCACACCGTCCTCATCTCCACCCAGCATGATGAGACCGTCACCAACGATGAGATCGCTG	682
SEQ	ID	No:	42	701	CGACCTCAAGGAGCATGTCAACAAGCCGGTGATCCCCGCAAAAGTAGCTCGATGAGAACA	760
SEQ	ID	No:	43	683	CAGACCTCAAGGAGCATGTCAACAAGCCGGTGATCCCCGGAAGTAGCTCGATGAGAACA	742
SEQ	ID	No:	42	761	CCATCTTCCACCTGAACCCCGTCTGGCCGCTTCGTATCGGGGGCCCCACGGTGACGCCG	820
SEQ	ID	No:	43	743	CCATCTTCCACCTGAACCCCATCGGGCCGCTTGTATCGGTGGCCCTCACGGCGATGCCG	802
SEQ	ID	No:	42	821	GTCTCACCGGCGCAAGATCATCATGACACCTATGGTGGCTGGGGAGCCACACGGCGCGG	880
SEQ	ID	No:	43	803	GTCTCACCGGCGCAAGATCATCATGACACCTATGGTGGTGGGGAGCCACACGGCGCGG	862
SEQ	ID	No:	42	881	GTGCCTTCTCTGGCAAGGACCCCAACRAAGGTCGACCCYAGTGGCGCCTACATTGCCAGGC	940
SEQ	ID	No:	43	863	GTGCCTTCTCTGGCAAGGACCCCTACCAAGGTCGACCGCGAGTGGCGCCTACATTGCCAGGC	922

